

# LitCovid track Multi-label topic classification for COVID-19 literature annotation

## README

Last updated on 08/31/2021

### Task description

The LitCovid track calls for a community effort to address automated topic annotation for COVID-19 literature. Topic annotation in LitCovid is a multi-label document classification task that assigns one or more labels to each article. There are 7 topic labels used in LitCovid: Treatment, Diagnosis, Prevention, Mechanism, Transmission, Epidemic Forecasting, and Case Report. These topics have been demonstrated to be effective for information retrieval and have also been used in many downstream applications related to COVID-19. However, annotating these topics manually has been a significant curation bottleneck. Increasing the accuracy of automated topic prediction in COVID-19-related literature would be a timely improvement, beneficial to curators and researchers worldwide.

### References

- Chen, Q., Allot, A. and Lu, Z., 2020. [Keep up with the latest coronavirus research](#). Nature, 579(7798), pp.193-193.
- Chen, Q., Allot, A. and Lu, Z., 2021. [LitCovid: an open database of COVID-19 literature](#). Nucleic Acids Research, 49(D1), pp. D1534-D1540.
- Add your review paper.

### Datasets

The training and development datasets contain 24,960 and 6,239 articles from LitCovid, respectively. The topics of the articles in both datasets have been manually reviewed.

The training and development datasets are provided in csv format, with the following fields retrieved from PubMed/LitCovid:

- pmid: PubMed Identifier
- journal: journal name
- title: article title
- abstract: article abstract
- keywords: author-provided keywords
- pub\_type: article type, e.g., journal article
- authors: author names
- doi: Digital Object Identifier
- label: annotated topics, i.e., **the output**
  - each article can be assigned one or more labels (Treatment, Diagnosis, Prevention, Mechanism, Transmission, Epidemic Forecasting, and Case Report)
  - each label is separated by a semicolon, e.g., 'Diagnosis;Treatment' means that the article is assigned both the label Diagnosis and the label Treatment

Other fields are also available via <https://ftp.ncbi.nlm.nih.gov/pub/lu/LitCovid/>, which provides additional fields if needed, such as biological entity annotations.

### Test set (updated in 08/22/2021)

The test dataset has been released with the same format as above, except the topic labels should be predicted by the participants. Submissions will be evaluated using both label-based and instance-based metrics that are commonly applied for multi-label classification. Evaluation scripts

are provided via [https://github.com/ncbi/biocreative\\_litcovid](https://github.com/ncbi/biocreative_litcovid). An example of prediction file is also provided via [https://github.com/ncbi/biocreative\\_litcovid/blob/main/prediction\\_label\\_samples.csv](https://github.com/ncbi/biocreative_litcovid/blob/main/prediction_label_samples.csv). The submission instruction will be available later.

### Submission instructions (updated in 08/31/21)

- Each team may submit up to five runs (predictions).
- Submissions are due 12<sup>th</sup> September 11:59:59 “Anywhere on Earth”.
- The prediction files must follow the same format as [https://github.com/ncbi/biocreative\\_litcovid/blob/main/prediction\\_label\\_samples.csv](https://github.com/ncbi/biocreative_litcovid/blob/main/prediction_label_samples.csv) and the evaluation script in [https://github.com/ncbi/biocreative\\_litcovid](https://github.com/ncbi/biocreative_litcovid) also validates the file format.
- Please submit each prediction separately with specific naming format (see the steps below).
- Submission steps:
  1. Please go to <https://easychair.org/conferences/?conf=bc7>
  2. Select ‘Track 5- LitCovid track Multi-label topic classification for COVID-19 literature annotation’ and click ‘Continue’, which will lead to the submission page
  3. In the submission page:
    - Please fill in the details of team members (authors) as requested
    - In the **title** field, please follow the format of *team\_name-submission\_id*, e.g., WorldPeaceTeam\_Submission1
    - In the **abstract** field, please provide a brief summary of the methods and models for this specific submission (up to one paragraph)
    - Please provide keywords as requested
    - File upload:
      - Please place your submission file into a folder and zip it (the submission portal requires the submission file type as zip)
      - Optionally, you could provide additional readme files into the folder if needed
      - Upload the zipped folder
    - Click ‘Submit’
    - Repeat the steps for another submission (up to five)

### Contract

Please contact [qingyu.chen@nih.gov](mailto:qingyu.chen@nih.gov) with the subject heading "BioCreative Track 5 LitCovid questions" if you have any questions.

### Status updates and FAQs

We will also provide updates and FAQs via

<https://biocreative.bioinformatics.udel.edu/tasks/biocreative-vii/track-5/>

